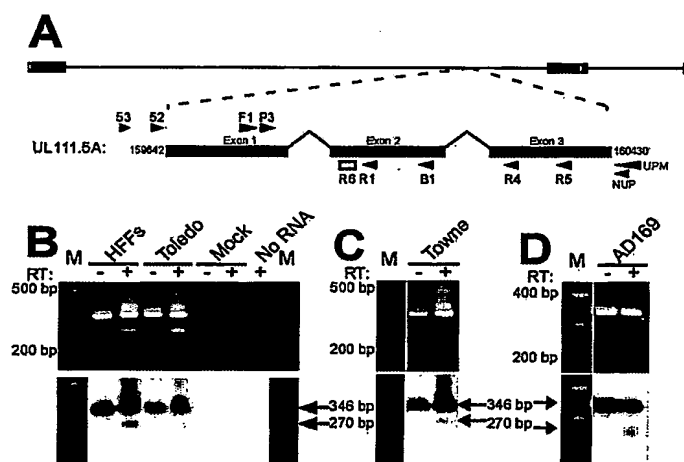
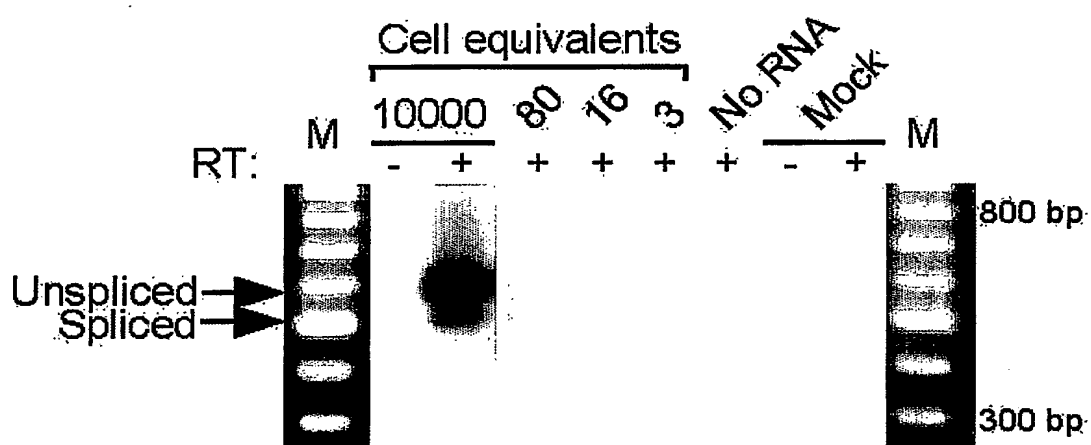
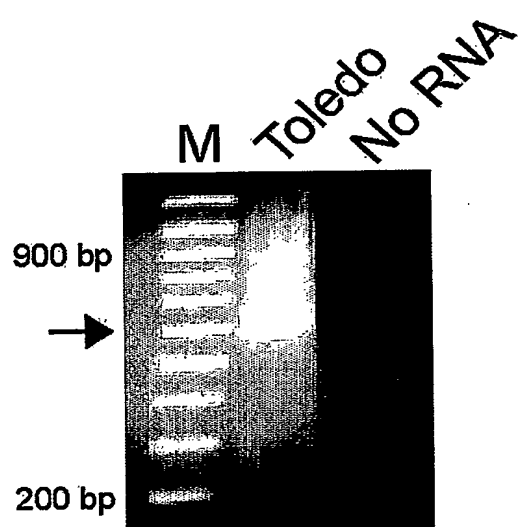


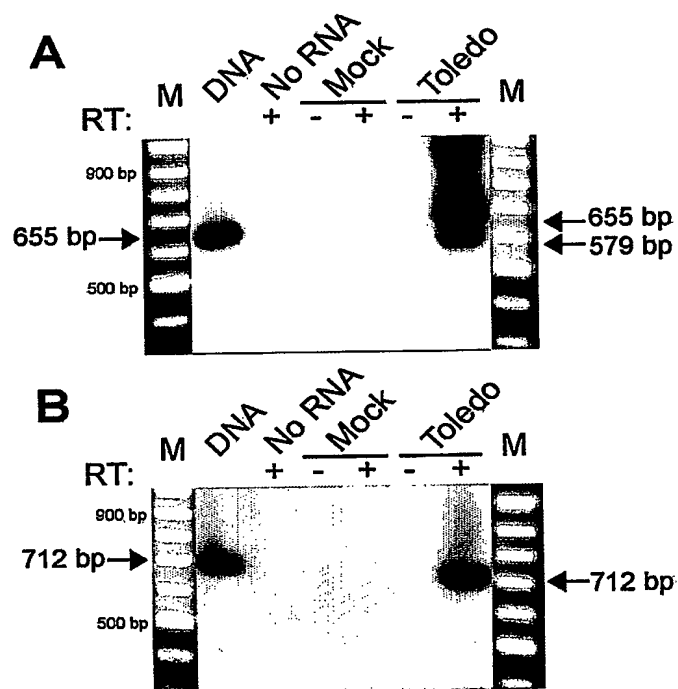
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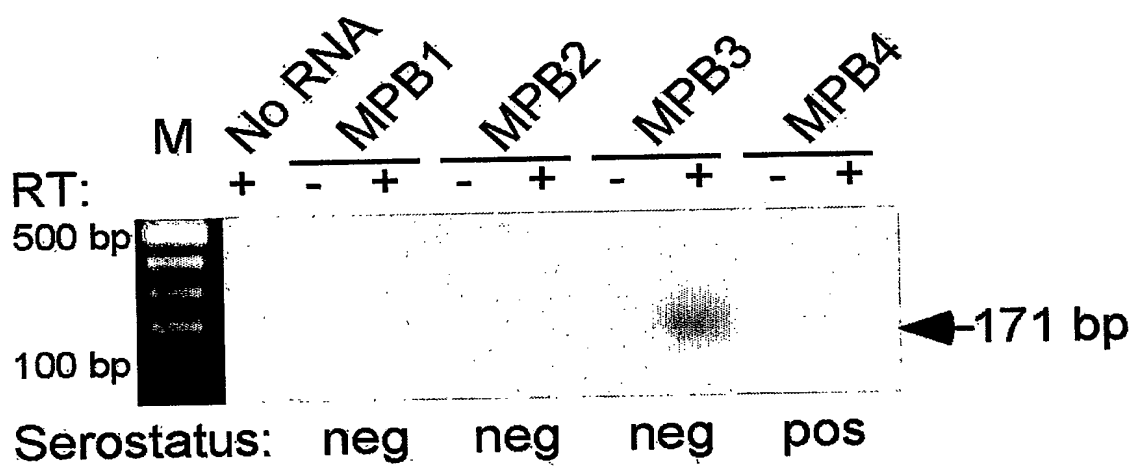




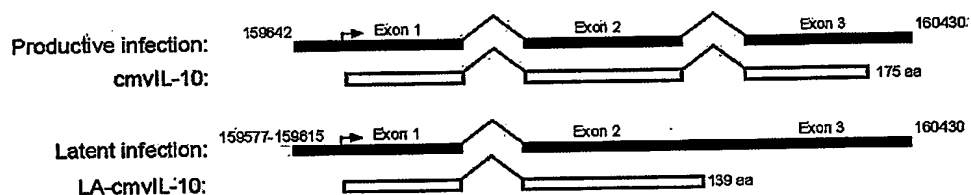
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A



B

Human IL-10: MHSSALLCCLVLLTGVRASPGGTQSENSCTHFPGNLRNMLRDLRDAFSSRVKTFFO 56
LA-cmvIL-10: MLSVMVSSSLVLIIVFFLGASEEAKPATTTIKNTKPKQCRPEDYATR LQDLRVTFHRVKPTLQ 61
cmvIL-10: MLSVMVSSSLVLIIVFFLGASEEAKPATTTIKNTKPKQCRPEDYATR LQDLRVTFHRVKPTLQ 61

Human IL-10: MKDQLDNLLLKESLLEDFKGYLGCCQALSEMIQFYLEEVMPPQAEHQDPPDIKAHVNSLGENLK 117
LA-cmvIL-10: REDDY-SVWLDGTVV--KGCGWGCSSVMDWLLRRYLEIVFPAGDHVYPGLKTELHSMRSTLE 122
cmvIL-10: REDDY-SVWLDGTVV--KGCGWGCSSVMDWLLRRYLEIVFPAGDHVYPGLKTELHSMRSTLE 122

Human IL-10: TLRRLRRCHRF L P C E N K S K A V E Q V K N A F N K L Q E K G I Y K A M S E F D I F I N Y I E A Y M T M K I R N 178
LA-cmvIL-10: SIYKDMRQCQVSVSVAALSAQR 143
cmvIL-10: SIYKDMRQCPLLGCQKSVISRLSQEAERKSDNGTRKGLSELDTLFSRLEEYLHSRK 179

